

Sequence listing

Sequence No. : 1

Sequence length: 180

5 Sequence type: Nucleic acid

Strandedness: Double

Topology: Linear

Sequence kind: cDNA to mRNA

Sequence description

10 GGA AAG CTC CCT TTC TCA AGA ATG CCC ATC TGT GAA CAC ATG GTA GAG 48
 Gly Lys Leu Pro Phe Ser Arg Met Pro Ile Cys Glu His Met Val Glu
 1 5 10 15
 TCT CCA ACC TGT TCC CAG ATG TCC AAC CTG GTC TGC GGC ACT GAT GGG 96
 15 Ser Pro Thr Cys Ser Gln Met Ser Asn Leu Val Cys Gly Thr Asp Gly
 20 25 30
 CTC ACA TAT ACG AAT GAA TGC CAG CTC TGC TTG GCC CGG ATA AAA ACC 144
 Leu Thr Tyr Thr Asn Glu Cys Gln Leu Cys Leu Ala Arg Ile Lys Thr
 35 40 45
 20 AAA CAG GAC ATC CAG ATC ATG AAA GAT GGC AAA TGC 180
 Lys Gln Asp Ile Gln Ile Met Lys Asp Gly Lys Cys
 50 55 60

25 Sequence No. : 2

Sequence length: 398

Sequence type: Nucleic acid

Strandedness: Double

Topology: Linear

30 Sequence kind: cDNA to mRNA

Origin:

Animal name: Homo sapiens

Cell kind: Stomach cancer tissue

Clone name: HP00839

35 Sequence characteristics:

Characterization code: CDS

Existence position: 43..303

Sequence description

40 GCAGGCCCCA GCCAGCTCAG GCTACACTAT CCCAGGATCA GC ATG GCC GTC CGC 54

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